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Interaction between filaggrin mutations and neonatal cat exposure in atopic dermatitis

Letter to the editor

Jacob P. Thyssen¹, Tarunveer S. Ahluwalia², Lavinia Paternoster³, Natalia Ballardini^{4,5}, Anna Bergström⁴, Erik Melén^{4,5}, Bo Chawes², Jakob Stokholm², Jonathan O'B Hourihane⁶, Donnchadh M. O'Sullivan⁶, Peter Bager⁷, Mads Melbye^{7,8,9}, Mariona Bustamante^{10,11,12}, Maties Torrent^{12,13,14}, Ana Esplugues^{12,15,16}, Liesbeth Duijts^{17,18}, Chen Hu^{17,19,20}, Niels J. Elbert^{19,20}, Suzanne G.M.A. Pasmans²⁰, Tamar E.C. Nijsten^{17,18,20}, Andrea von Berg²¹, Marie Standl²², Tamara Schikowski²³, Gunda Herberth²⁴, Joachim Heinrich^{22,25,26}, Young-Ae Lee²⁷, Ingo Marenholz²⁷, Susanne Lau²⁸, John A. Curtin²⁹, Angela Simpson²⁹, Adnan Custovic³⁰, Craig E. Pennell³¹, Carol A. Wang³¹, Patrick G. Holt³², Hans Bisgaard^{2*}, Klaus Bønnelykke^{2*}

* These authors contributed equally to the study

- 1) Department of Dermatology and Allergy, Herlev and Gentofte Hospital, University of Copenhagen, Hellerup, Denmark.
- 2) COPSAC, Copenhagen Prospective Studies on Asthma in Childhood, Herlev and Gentofte Hospital, University of Copenhagen, Copenhagen, Denmark.
- 3) MRC Integrative Epidemiology Unit at the University of Bristol, Bristol, UK.
- 4) Institute of Environmental Medicine, Karolinska Institutet, Stockholm, Sweden
- 5) Sachs' Children and Youth Hospital, Södersjukhuset, Stockholm, Sweden
- 6) Paediatrics and Child Health and INFANT Centre, University College Cork, Ireland.
- 7) Statens Serum Institut, Copenhagen, Denmark.
- 8) Department of Clinical Medicine, University of Copenhagen.
- 9) Department of Medicine, Stanford University School of Medicine, Stanford, CA, USA
- 10) ISGlobal, Barcelona Institute for Global Health, Barcelona, Spain
- 11) Universitat Pompeu Fabra (UPF), Barcelona, Spain
- 12) CIBER de Epidemiología y Salud Pública (CIBERESP), Madrid, Spain.
- 13) Ib-salut, Area de Salut de Menorca, Spain
- 14) Institut d'Investigació Sanitària Illes Balears (IdISBa), Spain
- 15) FISABIO– Universitat Jaume I –Universitat de València Joint Research Unit of Epidemiology and Environmental Health. Avenida de Catalunya 21, 46020, Valencia, Spain
- 16) Nursing Department, Faculty of Nursing and Chiropody, Universitat de València, Av. Blasco Ibáñez, 13, 46010 Valencia, Spain
- 17) Department of Pediatrics, Division of Respiratory Medicine and Allergology, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands.
- 18) Department of Pediatrics, division of Neonatology, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands.
- 19) The Generation R Study Group, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands.
- 20) Department of Dermatology, Erasmus MC, University Medical Center Rotterdam, Rotterdam, The Netherlands. Research Institute, Department of Pediatrics, Marien-Hospital Wesel, Wesel, Germany
- 21) Research Institute, Department of Pediatrics, Marien-Hospital Wesel, Wesel, Germany
- 22) Institute of Epidemiology, Helmholtz Zentrum München - German Research Center for Environmental Health, Neuherberg, Germany
- 23) IUF-Leibniz Research Institute for Environmental Medicine, Düsseldorf, Germany
- 24) Department of Environmental Immunology/Core Facility Studies, Helmholtz Centre for Environmental Research – UFZ, Leipzig, Germany
- 25) Institute and Clinic for Occupational, Social and Environmental Medicine, University Hospital, Ludwig Maximilian University (LMU), Munich, Germany

- 26) Allergy and Lung Health Unit, Melbourne School of Population and Global Health, The University of Melbourne, Melbourne, Australia.
- 27) Max-Delbrück-Center (MDC) for Molecular Medicine, Berlin, Germany; Clinic for Pediatric Allergy, Experimental and Clinical Research Center of MDC and Charité Universitätsmedizin Berlin, Berlin, Germany
- 28) Department of Pediatrics, Division of Pneumology, Immunology Charité, Universitätsmedizin Berlin, Berlin, Germany
- 29) Division of Infection Immunity and Respiratory Medicine, School of Biological Sciences, The University of Manchester, Manchester Academic Health Science Centre, and Manchester University NHS Foundation Trust
- 30) Section of Paediatrics, Department of Medicine, Imperial College London, London, United Kingdom
- 31) School of Medicine and Public Health, Faculty of Medicine and Health, University of Newcastle, Callaghan, New South Wales, Australia
- 32) Telethon Kids Institute, University of Western Australia, Nedlands, Australia

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Correspondence:

Professor Klaus Bønnelykke, MD, PhD

COPSAC, Copenhagen Prospective Studies on Asthma in Childhood

Herlev and Gentofte Hospital

Ledreborg Alle 34

2820 Gentofte

Denmark

E-mail: kb@copsac.com

Website: www.copsac.com

Introduction

Atopic dermatitis (AD) is a prevalent inflammatory skin disease. Loss-of-function mutations in filaggrin gene (*FLG*) represent the strongest genetic risk factors for AD, being strongly associated with early disease onset and persistence into adulthood.¹ The epidermis of individuals with mutations in *FLG* is fundamentally different from normal skin being characterized by increased penetration of allergens.²

Recent birth cohort studies showed a significant interaction between cat ownership at birth and mutations in *FLG* (R501X, 2282del4) on the development of early-onset AD.³ This finding was replicated for the 2282del4 *FLG* mutation in a Dutch cohort study, and extended to further associate with risk of allergic sensitization.⁴ We performed analyses in multiple birth cohorts to examine the consistency and overall strength of the previously observed interaction.

Materials and methods

Cohorts, exposures, genotypes and phenotypes

Consortium collaborators were invited to participate in the study,⁵ and 13 birth cohorts provided data on cat exposure, AD, and filaggrin mutations (Table 1 and Supplementary Table 1+2). All cohorts had information on the most common mutations in *FLG*, R501X and 2282del4, and the majority also had information on R2447X and S3247X (Table 1). Heterozygous, compound heterozygous and homozygous *FLG* mutation carriers were pooled as mutation carriers. Cat ownership/exposure was based on questionnaires or interviews. AD diagnoses were based on questionnaires in 11 cohorts, and by physician examination in 3 cohorts (COPSAC2000, COPSAC2010 and MAAS). For further details, please see supplementary information online.

Statistical analysis and outcomes

The predetermined primary outcome was AD onset before one year of age (AD_{early}). Secondary outcomes included i) current AD at seven years of age or the year of assessment closest to, but before, 7 years ($AD_{current}$), and ii) a history of AD during the first 7 years of life, or last year of assessment (AD_{ever}). For further details, please see supplementary information online.

Results

A total of 22,133 children were studied (Table 1 and supplementary Table 1). The median prevalence (range) of mutations in *FLG* was 9.4% (4.6-12.3), cat exposure 15% (7.9-29.6), AD_{early}, AD_{current}, and AD_{ever}, respectively, 18% (9.7-34.6), 13.9% (3.9-20), and 39.5% (20.4-67). There was no interaction between *FLG* mutations and cat exposure on the risk of the primary outcome ‘AD_{early}’ (OR 1.10 (95% CI 0.86-1.43), I²% 0.0), (Figure 1 and Table 1). There was a statistically significant interaction for the secondary outcome of having AD at last time of examination or questioning at 7 years of age (AD_{current}), in the direction of increased risk of AD from cat exposure in children with *FLG* mutations (OR 1.36 (95% CI 1.02-1.82) I²% 8.6), but this was not statistically significant after adjustment for multiple testing. The *FLG*-stratified analyses showed a trend towards cat exposure being a risk factor in children with *FLG* mutations and a protective factor in children without *FLG* mutations (Figure 1 and Supplementary Table 3). No interaction was found for the other secondary outcome ‘AD_{ever}’ (OR 1.06 (95% CI 0.82-1.37), I²% 0.0)

Discussion

We found no interaction between cat exposure in infancy and mutations in *FLG* on ‘early-onset AD’ or ‘AD ever’. A nominally significant interaction in the expected direction was found for the secondary outcome ‘current AD’ at 7 years of age, but this did not survive adjustment for multiple testing.

A particular study strength is the large number of independent birth cohorts with prospective assessment of exposure and outcomes. Most cohorts had genotype information for the 4 most common *FLG* mutations ensuring a high degree of correct classification. AD diagnoses were based on questionnaire data in most cohorts, potentially reducing diagnostic specificity. Since AD is a chronic and relapsing disease, short episodes of other eczemas, e.g. due to irritant or allergic contact dermatitis, may be suspected of being AD by parents and caregivers, in particular in the first years of life where flexural accentuation is not yet occurring. One may argue that AD measured at 7 years of age is expected to have a higher specificity due to flexural involvement.⁶ Notably, the high prevalence of early AD in some cohorts could mask a true cat exposure-*FLG* mutation

interaction. Cat exposure was only assessed around birth, and it is possible that later exposure to cat could have an unmeasured effect on AD. Similar, the extent of cat exposure might vary between studies and families. Other environmental factors were not included since covariate availability differed between the cohorts. Reverse causality cannot be excluded, as families who had experienced atopic disease might have avoided having pets to prevent allergic disease in their (next) child. However, one would expect families with *FLG* mutations, and thereby increased risk of eczema, to avoid cat ownership, which would tend towards an apparent protective effect of having a cat.

No association between cat ownership and AD was found in another meta-analysis of 13 studies (relative risk 0.94 (95%CI 0.76-1.16)).⁷ When a compelling gene-environment interaction was observed between *FLG* mutations and cat ownership on the risk of early-onset AD in birth cohorts, it raised the possibility that preventive measures against pediatric AD could be identified by taking the genetic susceptibility into account.³
⁴ The COPSAC2000 study, which provided the basis for the previous report of interaction between cat and *FLG* mutations,³ benefited from close follow-up of children, and high AD diagnostic accuracy, whereas most birth cohorts in the present meta-analysis used questionnaires, potentially leading to misclassification.

No pathomechanism has been established for the proposed association between cat ownership and AD in *FLG* mutation carriers. Possibly, very small cat allergens might penetrate into the viable layers of the epidermis, where they can exert immune effects.⁸ Previous studies demonstrating increased risk of peanut allergy *FLG* mutation carriers, also suggest increased peanut allergen skin penetration.⁹ Another explanation could be an effect of cat exposure on the gut microbiome of mothers and children. However, the absence of filaggrin protein expression in the gut argues against this.

In conclusion, this meta-analysis could not confirm an interaction between cat exposure in infancy and *FLG* mutations on development of early-onset AD. Gene-environment interactions remain largely unknown.

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Table 1. Baseline characteristics of participants in birth cohorts.

| Cohort | ALSPAC | BAMSE | Baseline | COPSAC 2000 | COPSAC 2010 | DNBC | Generation R | GINplus | INMA* | LISA | MAS | MAAS | RAINE |
|--|--------------------------------------|---|--------------------------------------|--|--------------------------------------|--------------------------------------|--------------------------------------|--------------------------------------|--------------------------------------|--|---|--------------------------------------|--------------------------------------|
| Cohort inclusion criteria | Children from the general population | Children from the general population | Children from the general population | Children born from mothers with asthma | Children from the general population | Children from the general population | Children from the general population | Children from the general population | Children from the general population | Children from the general population | Children from the general population | Children from the general population | Children from the general population |
| Study year baseline | 1991-1992 | 1994-1996 | 2008 | 2000 | 2010 | 1996-2001 | 2002-2006 | 1995-98 | 1997-2006** | 1997-99 | 1990 | 1996-1997 | 1989-1991 |
| Filaggrin mutations genotyped | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4, R2447X | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4 | R501X, 2282del4, R2447X, S3247X | R501X, 2282del4 | R501X, 2282del4, R2447X, S3247X | R501X; 2282del4, R3247X, R2447X | R501X, 2282del4, S3247X, rs138726443 |
| Proportion with FLG mutations | 11% (834/7743) | 7.2% (138/1906) | 10.8% (146/1344) | 12.3% (49/396) | 10.3% (72/700) | 9.7% (91/935) | 9.4% (268/2849) | 7.0% (104/1490) | 4.6% (28/606) | 7.0% (69/987) | 9.7% (79/813) | 10.1% (87/864) | 9.3% (140/1500) |
| Basis for atopic dermatitis diagnosis | Questionnaire | Questionnaire | Questionnaire and clinical diagnosis | Clinical diagnosis | Clinical diagnosis | Questionnaire | Questionnaire | Questionnaire | Questionnaire | Questionnaire | Questionnaire and clinical diagnosis | Questionnaire | Questionnaire |
| AD 'early onset (≤1y)' | 18% (1368/7743) | 16.9% (323/1906) | 22.7% (292/1282) | 25.3% (100/396) | 11.1% (78/700) | 14.7% (137/935) | 21.6% (545/2521) | 11.3% (167/1477) | 31.7% (192/606) | 9.7% (94/973) | 14.14% (115/813) | 34.6% (160/462) | 22.8% (341/1498) |
| AD 'current' | 20% (1270/6402) | 17.2% (327/1896) | 15.2% (178/1168) | 13.9% (55/396) | 13.6% (95/700) | 6.8% (65/963) | 18.6% (496/2658) | 5.93% (77/1298) | 19.5% (118/604) | 3.9% (32/825) | 7.5% (58/773) | 14.1% (96/681) | 13.3% (184/1386) |
| AD 'ever (0-7 y)' | 67% (4367/6501) | 39.5% (743/1878) | 26.3% (354/1344) | 42.2% (175/396) | 27.6% (193/700) | 20.4% (196/963) | 41.4% (1180/2849) | 35.3% (447/1266) | 49.6% (307/618) | 32.6% (269/826) | 36.2% (294/813) | 60.7% (306/504) | 39.7% (596/1500) |
| AD assessment time-points | 6, 18, 30, 42, 57, 69, 81 months | 1, 2, 4 and 8 years | 6, 12 and 24 months | 1 month, and then every 6 months. | 1 month, and then every 6 months. | 6 and 18 month, and 7 years | 6 months, and 1, 2, 3, 4 and 6 years | 1, 2, 3, 4, 6 and 10 years | 1, 2, and 4 years | 6, 12, 18, 24 months and 4, 6 and 10 years | 1, 3, 6, 12, 18, 24 month and then yearly | 1,3,5,8 years | 1, 3, 5, 8 years |
| Child age at cat exposure assessment | During pregnancy | At baseline (3 months) and/or 1 year follow-up. | 6 months and 12 months | Birth | Birth | 18 months | age < 1 year) | 1 year | 1 year | 3 months and 1 year | 3 months | Birth | 1 year |
| Early life cat exposure % | 29.6 | 10.6** | 8.8 | 15 | 20 | 16.4-22.4 ** | 25.3 | 7.9 | 11.5 | 12.2 | 12.7 | 20.5 | 18 |

*INMA sub cohorts: VAL, SAB, MEN

** please see supplementary Table 1 for further details.

Figure 1. Interaction between cat exposure and common *FLG* mutations in relation to a) Early onset atopic dermatitis, b) Current atopic dermatitis and c) atopic dermatitis in the first 7 years of life.

